



Probing Uranium Resistance by the Aerobic Aquatic Bacterium *Caulobacter crescentus*

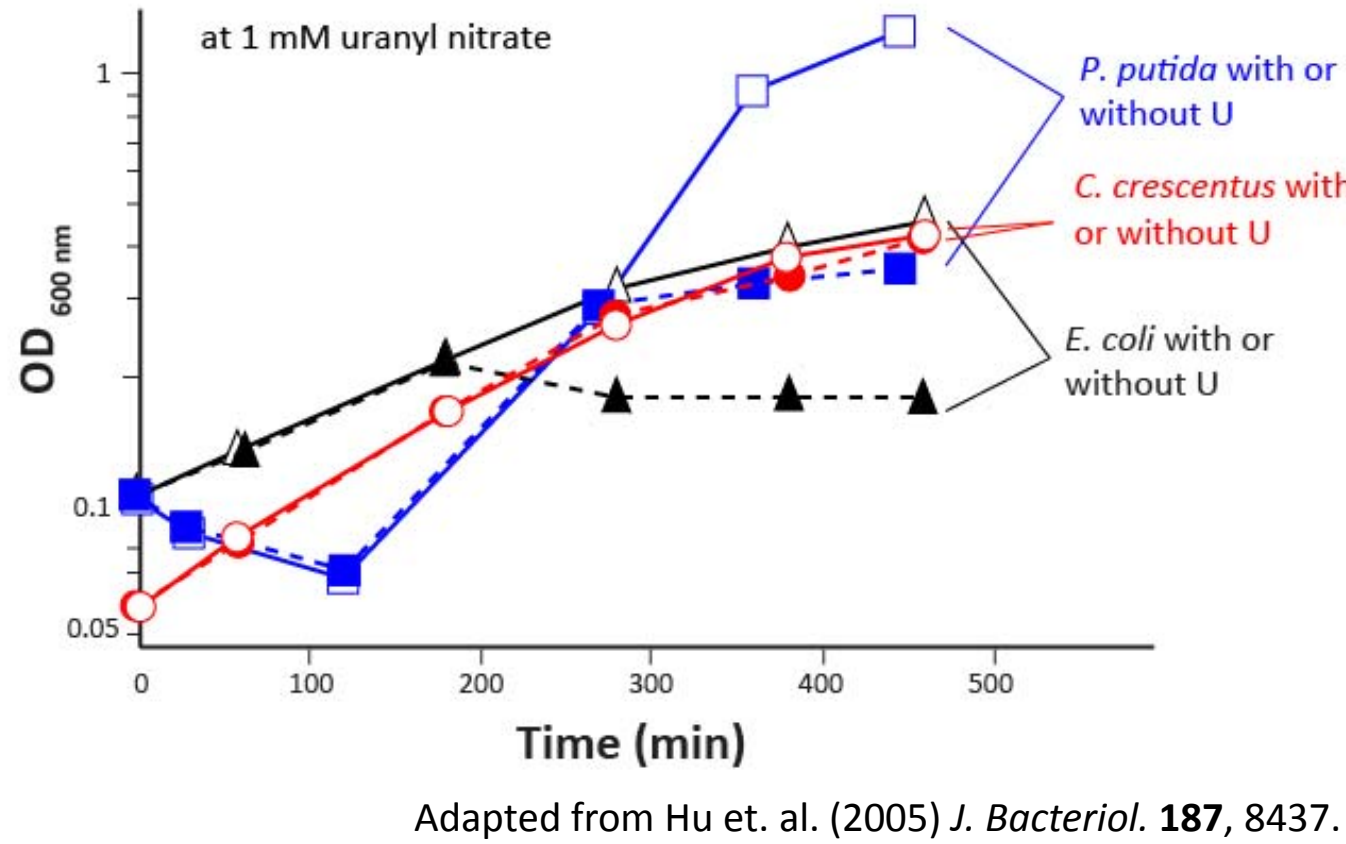
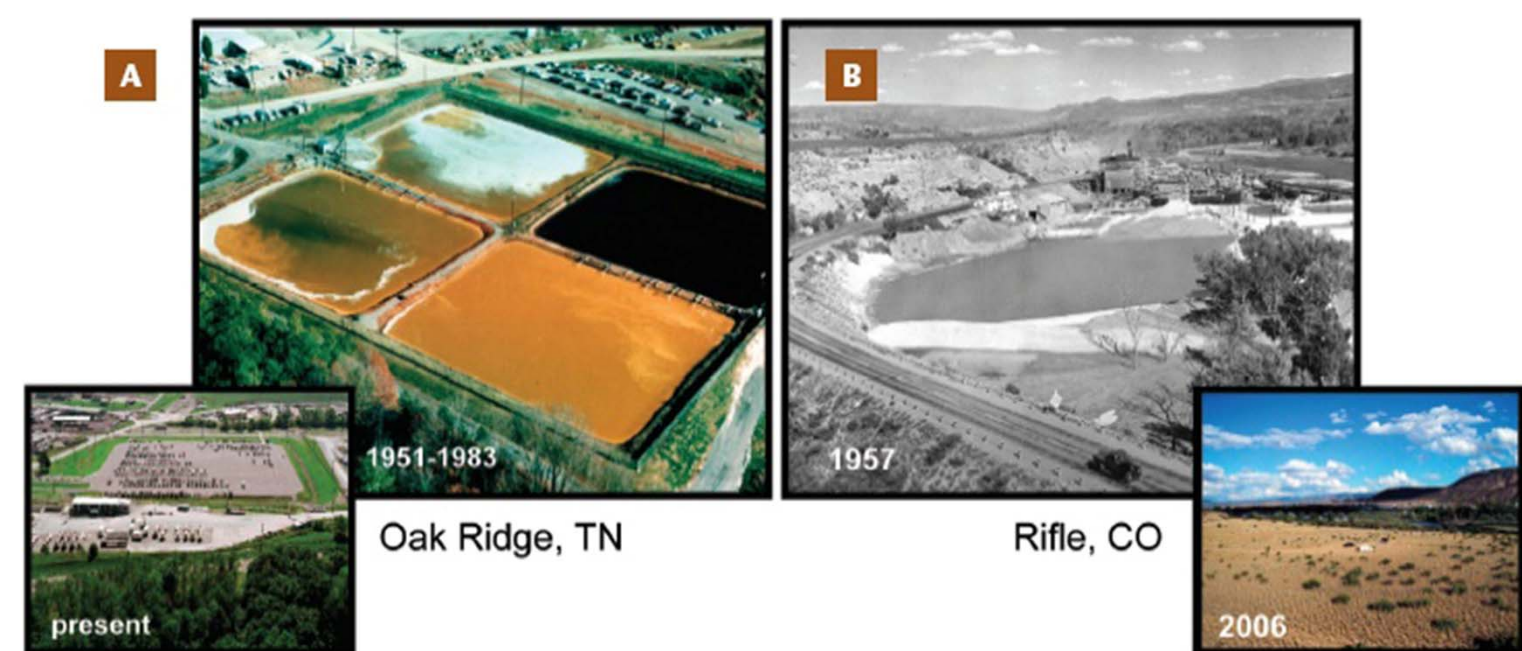


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C. crescentus is highly resistant to uranium:

- Uranium (U), particularly in its water soluble form U(VI), poses a significant threat to human health and wildlife as an environmental contaminant [1].
- The remediation of some 120 U-contaminated sites covering 7280 km² in the United States has been the focus and responsibility of the DOE. These sites have U(VI) levels up to 2000 times higher than the EPA regulated level of 0.03 mg/L (0.1 μM) [1].
- One strategy to help remediate U contamination is to use microbes which are highly tolerant of U(VI) and are able to mineralize U under aerobic conditions [2].
- The aerobic, aquatic, freshwater bacterium *Caulobacter crescentus* has been shown to be highly tolerant of U(VI). Our aim is to understand U detoxification and biomineralization processes in *C. crescentus* [3-4].



S-layer protective against U stress?:

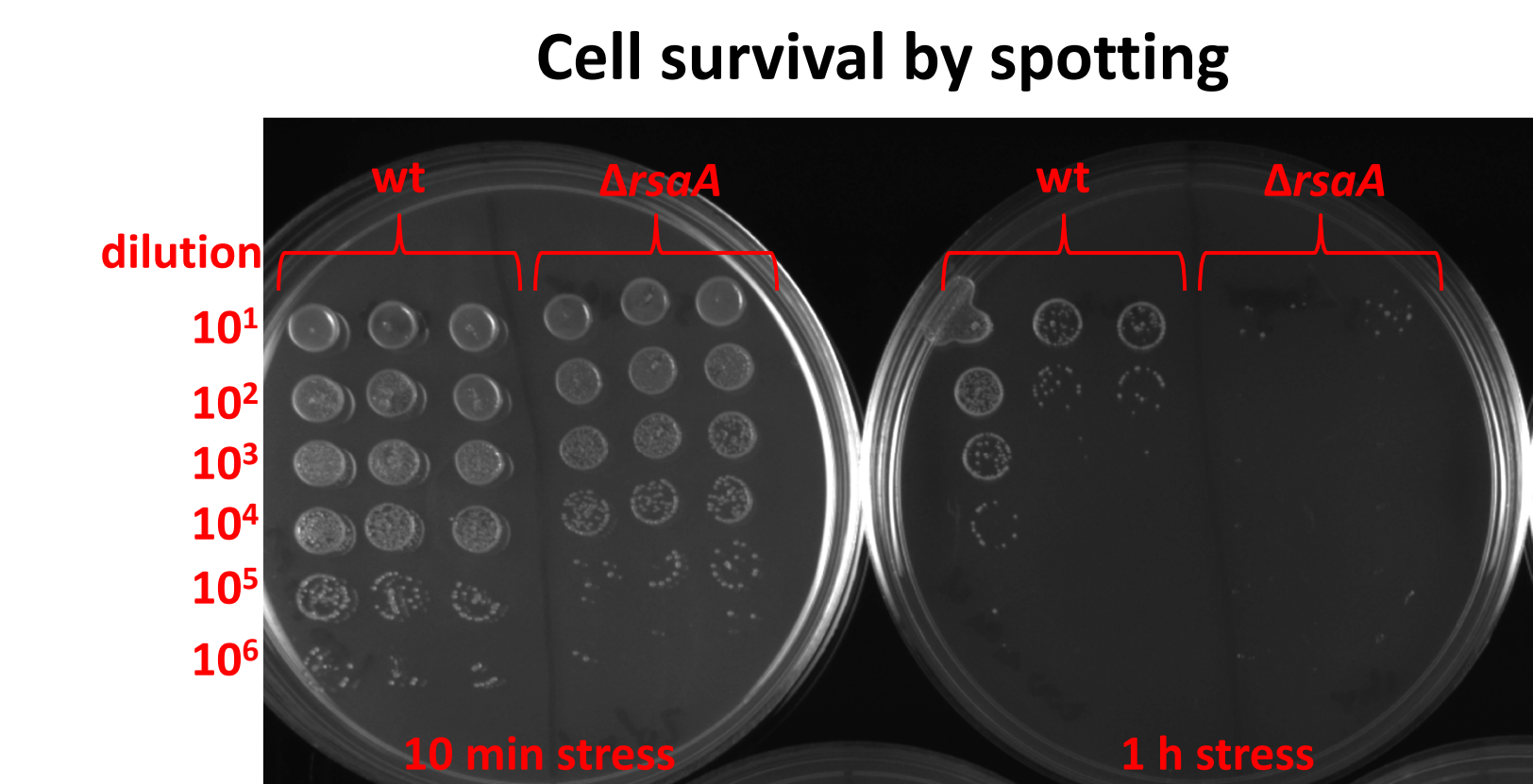
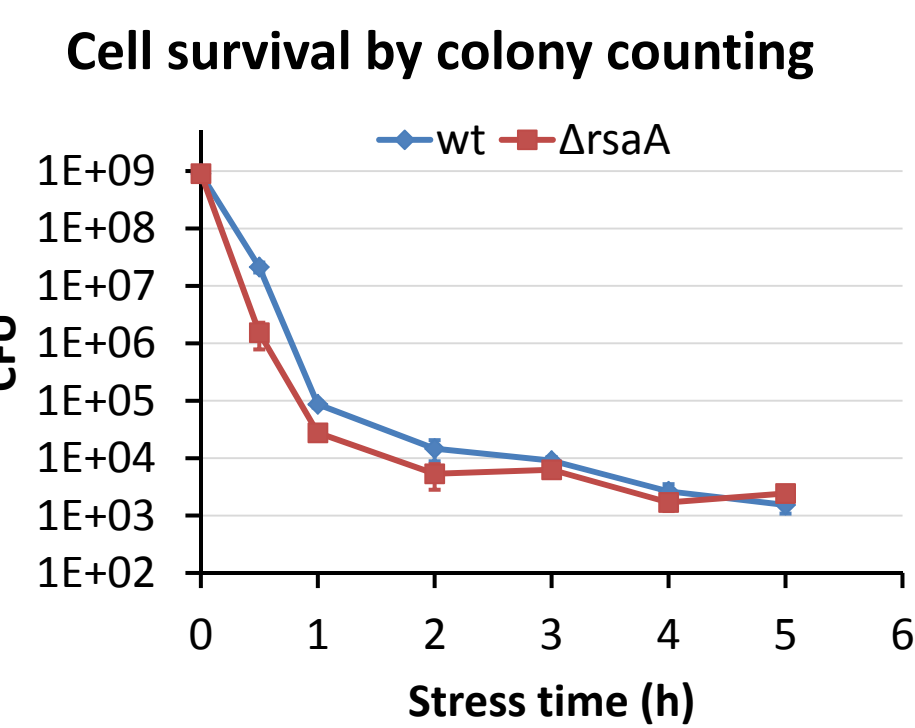
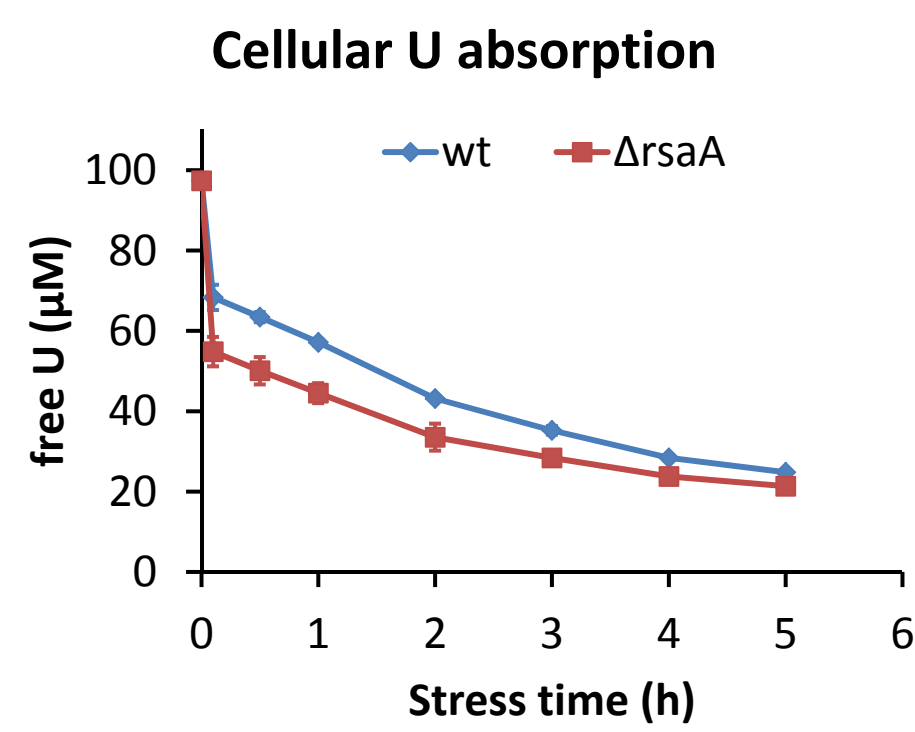
What is the S-layer?:

- The S-layer is the outermost layer of the cellular envelope for many bacteria and archaea.
- It consists of protein (RsaA in *C. crescentus*) or glycoprotein that self-assemble into an array on the bacterial surface [5].

S-layer mutant (Δ rsaA) is more susceptible to U compared to wt:

U stress challenge:

- Wt and Δ rsaA were grown in PYE medium to log phase and washed.
- Cells were re-suspended in 100 μM uranyl nitrate in 10 mM Tris pH 7.0 and incubated at 30 °C with rocking.
- CFU were determined by serial dilution followed by spotting/colony counting.
- The amount of U bound to the cells was determined using the ArsenazoIII assay.



Conclusions:

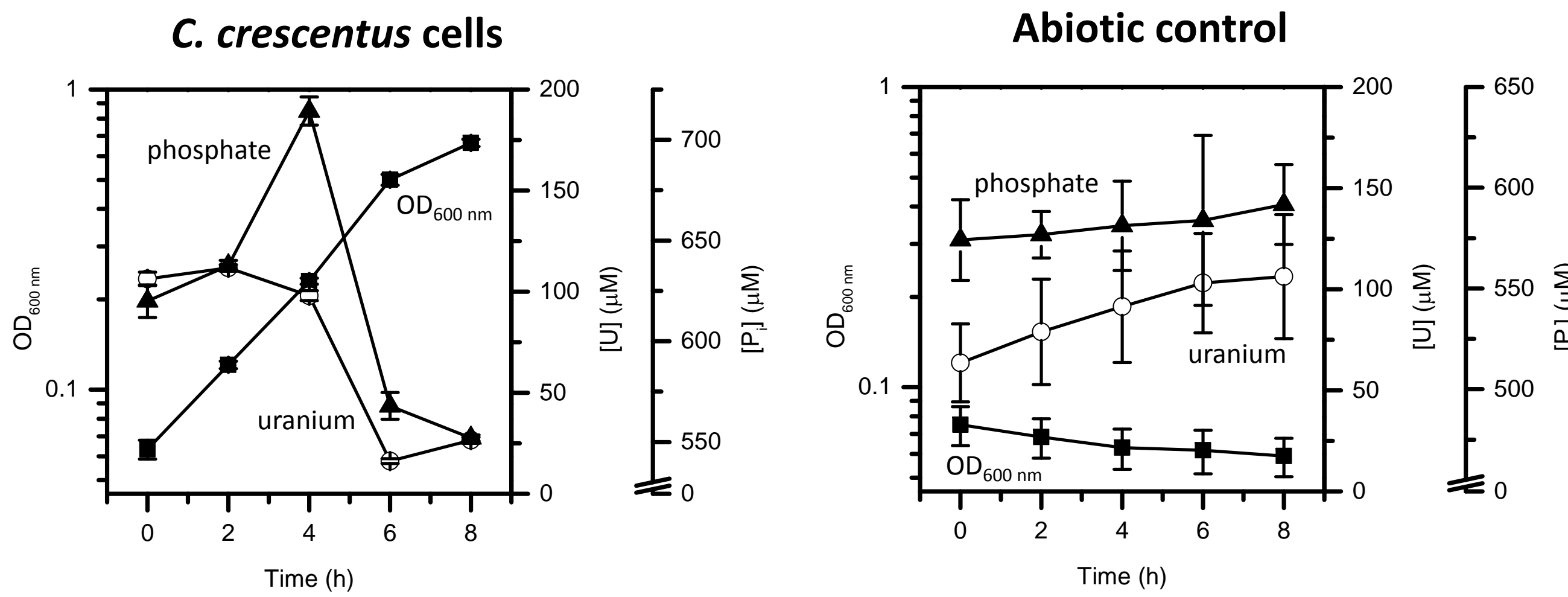
- Wt initially survives better than the S-layer mutant during the U stress challenge at pH 7, suggesting that the S-layer does have a somewhat protective role in U tolerance.
- Cellular U absorption is positively correlated with survival in the stress assay.

Future direction:

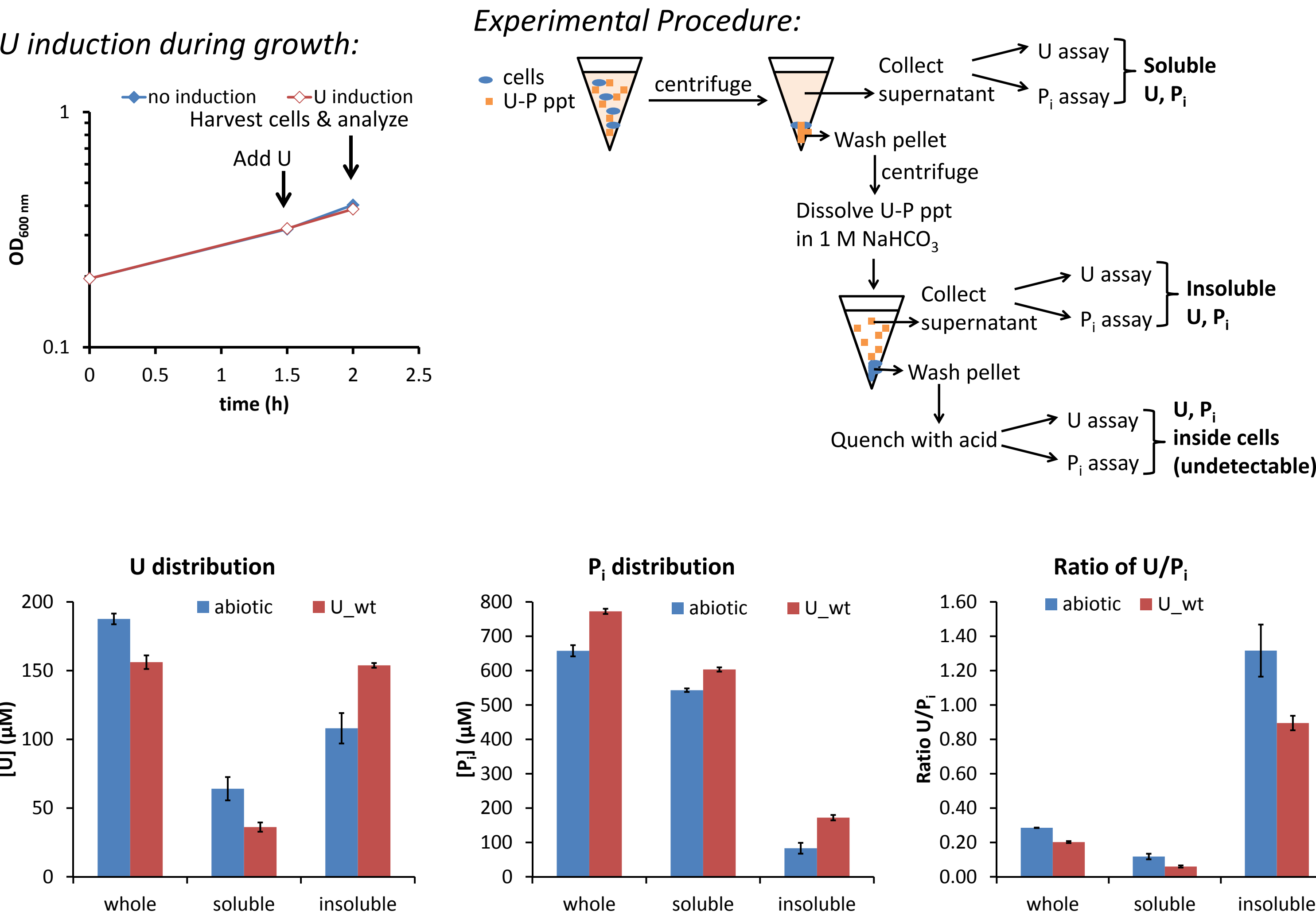
- Perform whole cell FTIR and TEM of the cells under the U stress challenge to determine if there is a difference in the U binding mode between mutant and wt.

Phosphate metabolism facilitates U precipitation:

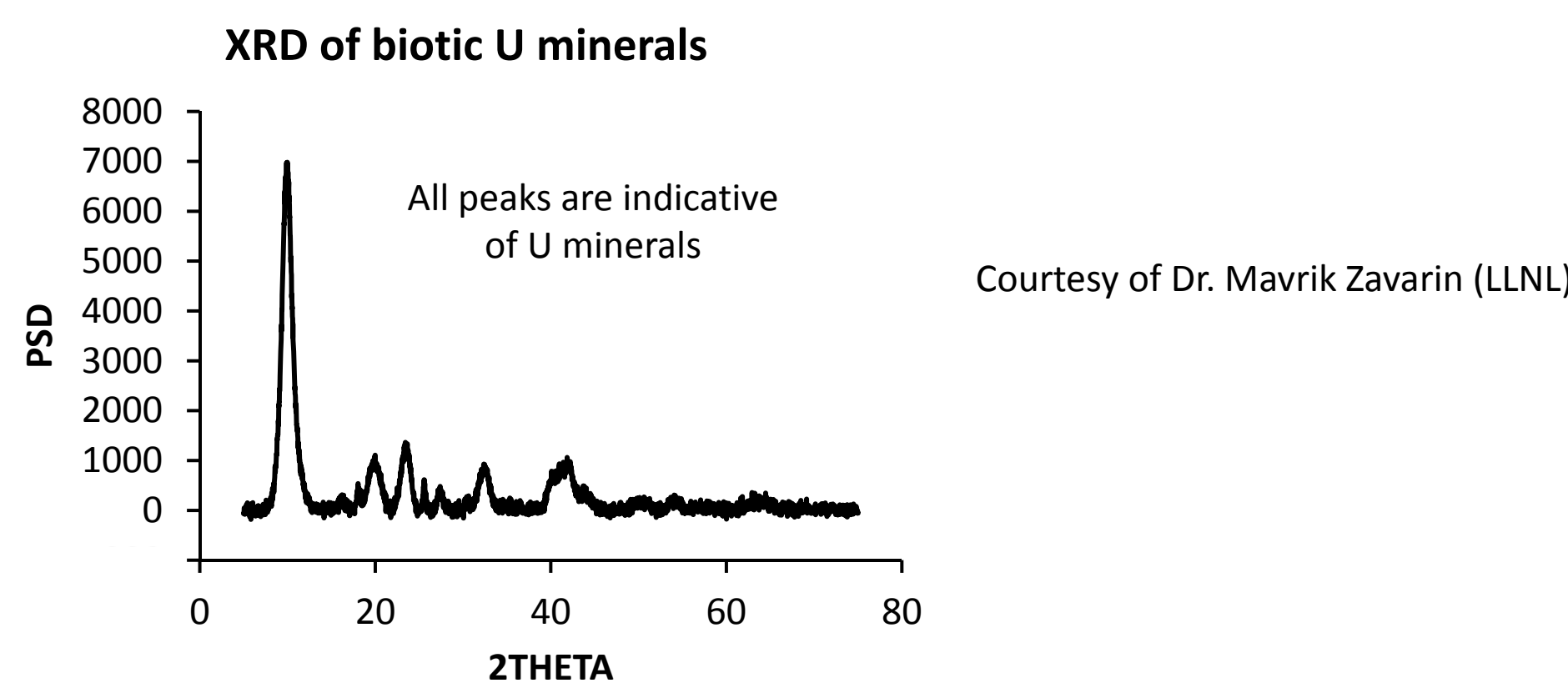
C. crescentus cells precipitate U and P_i concurrently during growth in PYE supplemented with 200 μM uranyl nitrate:



C. crescentus cells produce crystalline U-P minerals in higher amounts compared to the abiotic mineralization process:

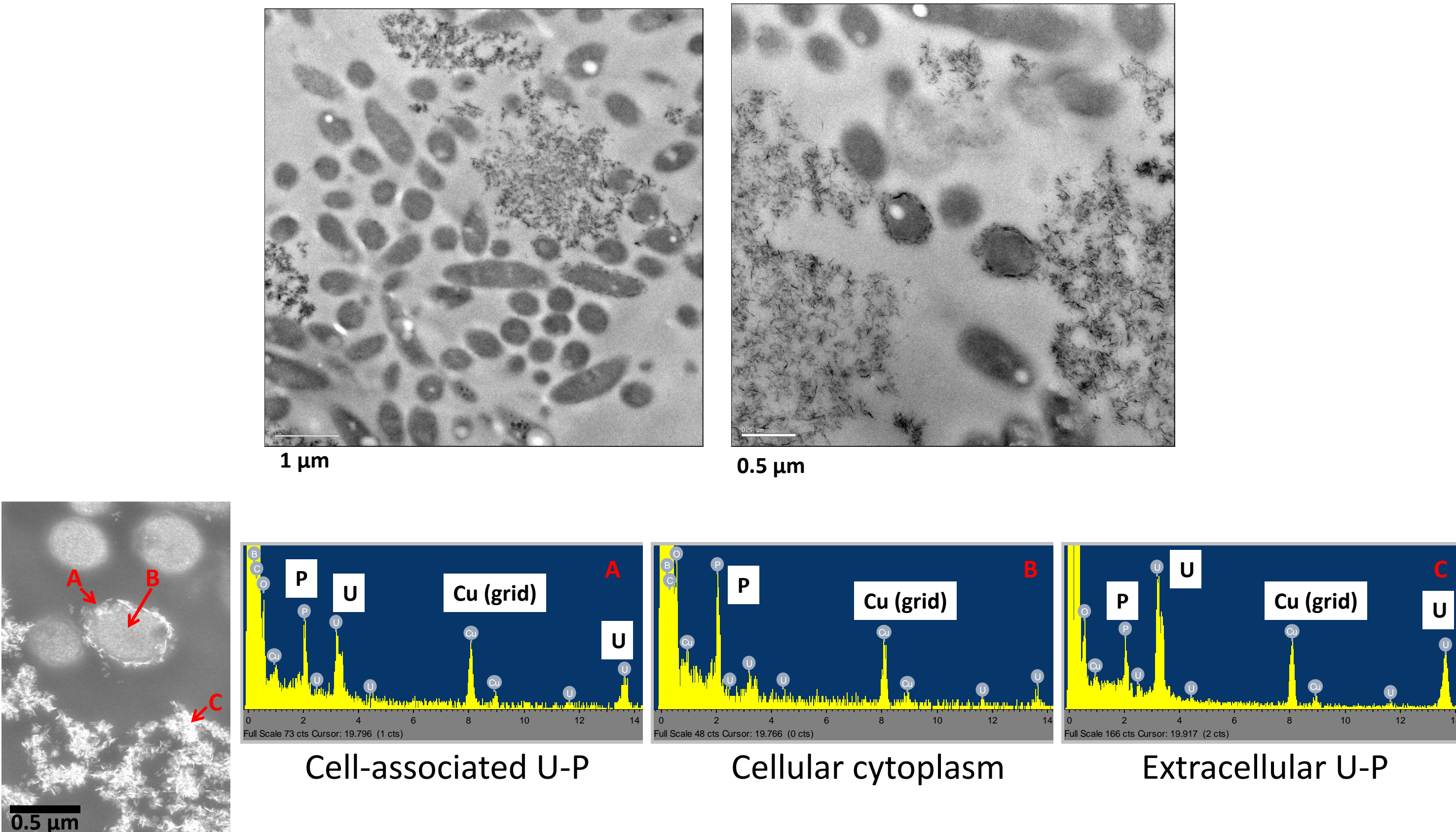


- Higher amounts of U-P minerals are produced biotically than abiotically.
- The ratio of U/P_i in the biotic minerals is lower than the abiotic minerals.



- Crystalline U-P minerals were not detected by XRD in an abiotic control or in a dead cell control.
- XRD analysis indicates the presence of:
(NH₄)(UO₂)(PO₄)·3H₂O (uramphite), K(UO₂)(PO₄)·3H₂O (potassium uranyl phosphate hydrate), and/or Na_{0.43}K_{0.57}(UO₂)(PO₄)·xH₂O (potassium sodium uranyl phosphate hydrate)

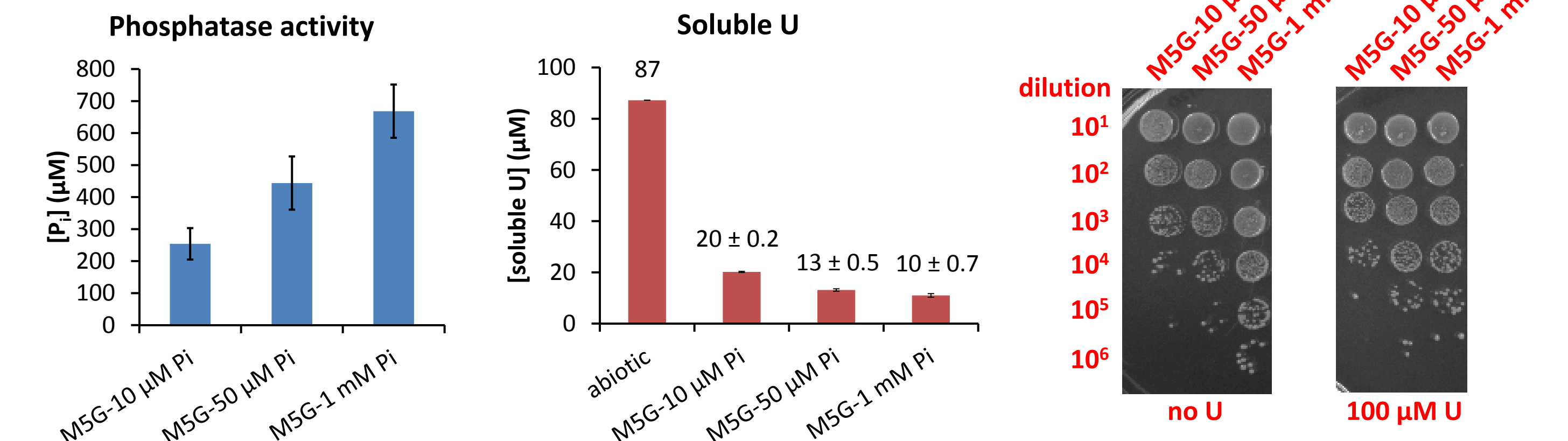
TEM coupled with EDS shows that the majority of the U is extracellular:



- No significant amount of U is found in the interior of the *C. crescentus* cells.

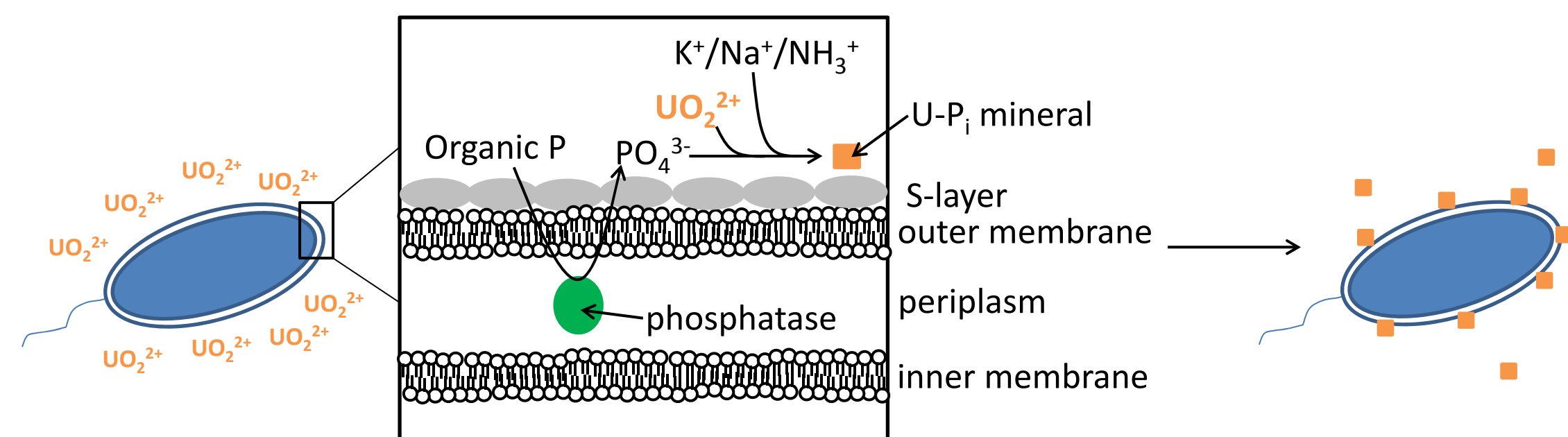
Cellular phosphatase activity is correlated with cell survival?

- C. crescentus* cells were grown in minimal medium (M5G) supplemented with 10 μM (very limited), 50 μM (limited), or 1 mM (excess) P_i.
- U stress challenge at 100 μM uranyl nitrate was performed in the presence of 5 mM β-glycerophosphate as an organic phosphate source for 2.5 h.



- There is a general growth defect with decreasing [P_i].
- There does not appear to be a significant correlation between level of phosphatase activity and survival at 50 and 100 μM U beyond the general growth defect.
- Future studies at higher [U] will be performed to determine if there is a correlation at higher [U].
- The presence of β-glycerophosphate in the U stress challenge improves survival. (Almost no colonies survive at 100 μM U without β-glycerophosphate).

Potential model for U defense in *Caulobacter crescentus*:



References:

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- NABIR Rep (2003), LLNL-42959.
- Hu P et. al. (2005) *J. Bacteriol.* **187**, 8437-8449.
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